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## CLAIMS

Equine herpesvirus (EHV) mutant, comprising one or more deletions, substitutions or insertions in the endogenous promoter region of an essential viral gene with respect to the parent strain.

- EHV mutant as claimed in claim-1, wherein deletions are introduced into the promoter.
- 3. EHV mutant as claimed in claims 1-2, wherein the gene is the Immediate Early gene.
- 4. EHV mutant as claimed in claims 1-3, wherein the mutant virus is the EHV-1 virus or the EHV-4 virus.
- 5. EHV mutant as claimed in claims 1-4, further comprising one or more mutations in one or more other genes and/or their promoters.
- 6. EHV-1 mutant as claimed in claims 1/5, comprising a deletion of the Saci-Saci fragment or the Hindili-Clai fragment or the Ndel-Ndel fragment or the Sphi-Sphi fragment of the promoter region of the Immediate Early gene.
- 7. Nucleic acid sequence, comprising the endogenous promoter region of an the immediate Early gene from EHV and optionally one or more flanking sequences, which promoter region comprises a deletion of the Sacl-Sacl fragment or the HindIII-Clal fragment or the Ndel-Ndel fragment or the Sphl-Sphl fragment of the promoter region of the Immediate Early gene.
- 8. Nucleic acid sequence as daimed in claim 7, wherein the EHV is EHV-1 or EHV-4.
- 9. Recombinant DNA molecule comprising a nucleic acid sequence as claimed in claims 7 or 8.
- 10. Host cell harbouring a recombinant DNA molecule as claimed in claim 9.
- 11. Vaccine comprising an EHV mutant as claimed in claims 1-6 and a pharmaceutically acceptable carrier or different.
- 12. A process for the preparation of an EHV mutant as claimed in claims 1-6, comprising transfection of a cell culture with a recombinant DNA molecule as claimed in claim 11 and EHV genomic DNA.
- 13. Method of genetically attenuating EHV, comprising mutation of the endogenous promoter region of an essential gene, which mutation consists of one or more deletions, substitutions/or insertions in the promoter region of an essential gene.
- 14. Method as claimed in claim, 13, wherein the EHV is EHV-1 or EHV-4.
- 15. Method as claimed in claims 13-14, wherein the gene is an Immediate Early gene.